



## COMPUTER SEARCH RESULTS

\*1: P58810. Lambda/chi-conoto...[gi:20454878] BLink, Links

>gi|20454878|sp|P58810|CXL4\_CONMR Lambda/chi-conotoxin MrIB (Chi-MrIB)  
VGVC CGYKLCHPC

\*2: P58808. Lambda-conotoxin ...[gi:20454876] BLink, Links

>gi|20454876|sp|P58808|CXL2\_CONMR Lambda-conotoxin CMrVIB precursor (Chi-conotoxin  
MrIA) (Chi-MrIA) (mr10a)  
MRCLPVLIIILLLTASAPGVVVLPKTEDDVPMSVYGNKGKSILRGILRNGVC CGYKLCHP  
C

\*3: P58807. Lambda-conotoxin ...[gi:20454875] BLink, Links

>gi|20454875|sp|P58807|CXL1\_CONMR Lambda-conotoxin CMrVIA  
VCCG YKLCHPC

4: P58809. Lambda-conotoxin ...[gi:20454877] Links

>gi|20454877|sp|P58809|CXL3\_CONMR Lambda-conotoxin CMrX  
GICCGVSFCYPC

5: AAG60408. conotoxin scaffold...[gi:12619493] BLink, Links

>gi|12619493|gb|AAG60408.1|AF214980\_1 conotoxin scaffold IX precursor [Conus pennaceus]  
MRCLPVFVILLLLTASGPSVDAKVHLKTKGDGPLSSFRDNAKSTLQRLQDKSTCCGYR  
MCVPCG

6: CAB95439. hypothetical prot...[gi:9366677] BLink, Links

>gi|9366677|emb|CAB95439.1| hypothetical protein [Trypanosoma brucei]  
MKFLVVDGDWPVTPYTLTRSELVSHSVEEADGGSEGGQLIIHGHTLVALCQQQLHTSVV  
RVRAAPDLSTLGLTILHESLSDSTGNIRLVTPKSLTRQVFAVDDLLVHGRLIVFTSRWESL  
ASAAIVGVKSDRREEHELTRRTKS RPEETKNSGRGNHDTSVSRPSVFSGRNRTALHKFTC  
AETYFQSIPPATVHDLDLLNNLIQAGTIRLDPRVEVYLCTVCGFLPRQTFTTSCCGAIMC  
GACTPLISTSSVMPSVMSEYVCIACKELPFTSHTAHVSRDAEVMKLVRELRLVHHPQL  
ASAASATSSHGMAEKSHASLPFLHSMGIMPAKRVDVGGAAPVLGSVFPNP

\*7: XP\_371536. similar to tripar...[gi:41124489] BLink, Domains, Links

>gi|41124489|ref|XP\_371536.1| similar to tripartite motif-containing 43 [Homo sapiens]  
MDSDFSHAFQKELTCVICLNLYLVDPTICCGHSFCRPCLCLSWEEAQSPANCPACREPS  
KMDFKTNILLKNLVTIARKASLWQFLSSEKQICGTHRQTKKMFCDDMDKSLCLLCSNSQ  
EHGAHKHYPIEEAAEEDREKLLKQMRILWKKIQENQRNLYEERRTAFLLRGDVVLRAQ  
MIRNEYRKLHPVLHKEEKQHLERLNKEYQEIFQQLQRSWVKMDQKSKHLKEMYQELM  
EMCHKPEVELLQDLGDIVARSESVLLHMPQPVNPELTAGPITGLVYRLNRFRVEISFHE  
VTNHNIRLFEDVRSWMFRGPNLSRSDYFAAWGARVFSFGKHYWELDVDNSCDWAL  
GVCNNSWIRKNSTMVNSEDIFLLCLKVDNHFNLTTSPVFPHYIEKPLGRVGVFLDFES  
GSVSFLNVTKSSLIWSYPAGSLTFPVRPFFYTGHR

\*8: NP\_620155. tripartite motif-...[gi:20270353] BLink, Domains, Links

>gi|20270353|ref|NP\_620155.1| tripartite motif-containing 43 [Homo sapiens]  
MDSDFSHAFQKELTCVICLNLYLVDPTICCGHSFCRPCLCLSWEEAQSPANCPACREPS  
KMDFKTNILLKNLVTIARKASLWQFLSSEKQICGTHRQTKKMFCDDMDKSLCLLCSNSQ  
EHGAHKHHPIEEAAEEHREKLLKQMRILWKKIQENQRNLYEEGRTAFLWRGNVVLRAQ  
MIRNEYRKLHPVLHKEEKQHLERLNKEYQEIFQQLQRSWVKMDQKSKHLKEMYQELM  
EMCHKPDVELLQDLGDIVARSESVLLHMPQPVNPELTAGPITGLVYRLNRFRVEISFHE  
VTNHNIRLFEDVRSWMFRGPNLSRSDYFAAWGARVFSFGKHYWELDVDNSCDWAL  
GVCNNSWIRKNSTMVNSEDIFLLCLKVDNHFNLTTSPVFPHYIEKPLGRVGVFLDFES  
GSVSFLNVTKSSLIWSYPAGSLTFPVRPFFYTGHR

9: NP\_041168. Zn-binding protei...[gi:9626898] BLink, Links

>gi|9626898|ref|NP\_041168.1| Zn-binding protein [Ictalurid herpesvirus 1]  
MELSLYRELPRKRRAVAQPRTRQPPKVHREDTGIVPGLRVESVVRPDSITNFRRLAAVK  
FKNPEVMECGGKGGCEECKTRPALMLNCDCTAKVPICLGCKPRAVRDHPCFNRNRTC  
SCGTIGVVAVVKLGCCGATLCDSCLGADWYPKSRTTRGTCGYCGAQTGLGVHYRPMKIIK  
NTRPYIPAETPIPKVQRHISARLLEEIRRCQLSRSEIVSLFGDHGSMMEYPTAVPAHLLAP  
KMIGYSDISKAFADDTLPRCSRAMCAANGVPGWRCGFQAPVLFPGMDMCLLCVVRQA  
EQTITQLVLNGTIWEDAPIAGTYHMSVWLDPTTVSVTPVDISAIDTFAVEFCGNLGSHHP  
EKFYRLSDLTCRIRWTAGGFLDPESLMVPEKERLKNTASAHSNNSH